

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/775,818

DATE: 03/05/2001
 TIME: 12:47:51

Input Set : A:\400684-SEQUENCE.txt
 Output Set: N:\CRF3\03052001\I775818.raw

3 <110> APPLICANT: Laboratory of Molecular Biophotonics
 5 <120> TITLE OF INVENTION: Method for selectively separating live cells expressing
 6 a specific gene
 8 <130> FILE REFERENCE: FP00-0043-00
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/775,818
 C--> 10 <141> CURRENT FILING DATE: 2001-02-05
 10 <150> PRIOR APPLICATION NUMBER: JP 2000/028117
 11 <151> PRIOR FILING DATE: 2000-02-04
 13 <150> PRIOR APPLICATION NUMBER: JP 2000/130793
 14 <151> PRIOR FILING DATE: 2000-04-28
 16 <160> NUMBER OF SEQ ID NOS: 20
 18 <170> SOFTWARE: PatentIn Ver. 2.1

**Does Not Comply
 Corrected Diskette Needed**

ERRORED SEQUENCES

248 <210> SEQ ID NO: 20
 249 <211> LENGTH: 15
 250 <212> TYPE: DNA
 251 <213> ORGANISM: Artificial Sequence
 253 <220> FEATURE:
 254 <223> OTHER INFORMATION: Probe
 256 <400> SEQUENCE: 20
 257 gaggttcctg tctgag
 E--> 260 FP00-0043-00
 E--> 261 1/8

15

→ Delete extraneous material from
 the end of the file. It is causing
 an invalid base count for sequence #20.

VERIFICATION SUMMARY

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:260 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:19 SEQ:20
L:260 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
M:254 Repeated in SeqNo=20
L:261 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:261 M:252 E: No. of Seq. differs, <211>LENGTH:Input:15 Found:20 SEQ:20